

# SEQUENCE LISTING

<110> Johal, Gurmukh S  
Multani, Dilbag S

<120> ISOLATED NUCLEIC ACID MOLECULES ENCODING THE DW3  
P-GLYCOPROTEIN OF SORGHUM AND METHODS OF MODIFYING  
GROWTH IN TRANSGENIC PLANTS THEREWITH

<130> 035718/275581

<140>

<141>

<150> 60/165,176

<151> 1999-11-12

<150> 09/711,619

<151> 2000-11-13

<160> 9

<170> PatentIn Ver. 2.1

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<211> 2139

<212> DNA

<213> Sorghum bicolor

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 <212> DNA  
 <213> Sorghum bicolor

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gcg	ttc	aac	gcg	gag	cgc	aag	atc	acg	ggg	ctg	ttc	gag	gcc	aac	ctg	192
Ala	Phe	Asn	Ala	Glu	Arg	Lys	Ile	Thr	Gly	Leu	Phe	Glu	Ala	Asn	Leu	
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Arg	Gly	Pro	Leu	Arg	Arg	Cys	Phe	Trp	Lys	Gly	Gln	Ile	Ala	Gly	Ser	
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Arg	Thr	Ile	Arg	Val	Phe	Met	Val	Leu	Met	Val	Ser	Ala	Asn	Gly	Ala	
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Asp	Asp	Val	Asp	Ala	Ala	Pro	Val	Pro	Glu	Arg	Pro	Lys	Gly	Glu	Val	
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Glu	Leu	Lys	His	Val	Asp	Phe	Ser	Tyr	Pro	Ser	Arg	Pro	Asp	Ile	Gln	
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gcg tac ggg cgc gag ggc gcg acg gag gcg gag gtg gtg gag gcg gcg	864
Ala Tyr Gly Arg Glu Gly Ala Thr Glu Ala Glu Val Val Glu Ala Ala	
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acg cag gcg aac gcg cac cgg ttc atc gcg gcg ctg ccg gag ggc tac	912
Thr Gln Ala Asn Ala His Arg Phe Ile Ala Ala Leu Pro Glu Gly Tyr	
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Gly Thr Gln Val Gly Glu Arg Gly Val Gln Leu Ser Gly Gly Gln Arg	
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Arg Thr Gly Trp Pro Arg Cys Ala Ala Arg Thr Pro Ser Arg Ser Ser	
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acc atc ccg acg ggt gct acg cgc gga tgc tgc agc tgc agc ggc	1245
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<211> 415

<212> PRT

<213> Sorghum bicolor

<400> 4

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 <213> Artificial Sequence

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 <223> Description of Artificial Sequence:oligonucleotide  
 primer designed from sequence of Zea mays Br2 gene

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<210> 6  
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Glu Leu Glu Ala Phe His Leu Pro Ser Pro Ala His Gln Pro Pro Gly	
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Phe His Leu Ala Ala Gly His Gln Pro Glu Ala Ala Ala Glu Gln Pro	
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Gly Ala Ala Pro Pro Ser Pro Ser Pro Pro Pro Pro Pro Ala Pro Leu	
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Glu Met Asp Gln Pro Pro Asn Ala Lys Pro Ala Ser Ser Ser Ala Ala	
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Ala Ala Gly Ala Asn Asp Asn Lys Lys Pro Thr Pro Pro Ala Ala Leu	
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Thr	Ile	Asp	Arg	Lys	Thr	Glu	Val	Glu	Pro	Asp	Asp	Val	Asp	Ala	Ala	
1105					1110				1115					1120		
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Pro	Val	Pro	Glu	Arg	Pro	Lys	Gly	Glu	Val	Glu	Leu	Lys	His	Val	Asp	

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Cys Gly Lys Ser Ser Val Leu Ala Leu Val Gln Arg Phe Tyr Glu Pro			
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Thr Ser Gly Arg Val Leu Leu Asp Gly Lys Asp Val Arg Lys Tyr Asn			
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ctg cgg gcg ctg cgg cgc gtg gtg gcg gtg gcg ccg cag gag ccg ttc			3648
Leu Arg Ala Leu Arg Arg Val Val Ala Val Ala Pro Gln Glu Pro Phe			
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ctg ttc gcg gcg agc atc cac gac aac atc gcg tac ggg cgc gag ggc			3696
Leu Phe Ala Ala Ser Ile His Asp Asn Ile Ala Tyr Gly Arg Glu Gly			
1220	1225	1230	
gcg acg gag gcg gag gtg gtg gag gcg gcg acg cag gcg aac gcg cac			3744
Ala Thr Glu Ala Glu Val Val Glu Ala Ala Thr Gln Ala Asn Ala His			
1235	1240	1245	
cgg ttc atc gcg gcg ctg ccg gag ggc tac ggg acg cag gtg ggc gag			3792
Arg Phe Ile Ala Ala Leu Pro Glu Gly Tyr Gly Thr Gln Val Gly Glu			
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cgc ggg gtg cag ctg tcg ggc ggg cag cgg cag cgg atc gcg atc gcg			3840
Arg Gly Val Gln Leu Ser Gly Gly Gln Arg Gln Arg Ile Ala Ile Ala			
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cgc gcg ctg gtg aag cag gcg gcc atc gtg ctg ctg gac gag gcg acc			3888
Arg Ala Leu Val Lys Gln Ala Ala Ile Val Leu Leu Asp Glu Ala Thr			
1285	1290	1295	
agc gcg ctg gac gcc gag tcg gag cgg tgc gtg cag gag gcg ctg gag			3936
Ser Ala Leu Asp Ala Glu Ser Glu Arg Cys Val Gln Glu Ala Leu Glu			
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cgc gcg ggg tcc ggg cgc acc acc atc gtg gtg gcg cac cgg ctg gcc			3984
Arg Ala Gly Ser Gly Arg Thr Thr Ile Val Val Ala His Arg Leu Ala			
1315	1320	1325	
acg gtg cgc ggc gcg cac acc atc gcg gtc atc gac gac ggc aag gtg			4032
Thr Val Arg Gly Ala His Thr Ile Ala Val Ile Asp Asp Gly Lys Val			
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gcg gag cag ggg tcg cac tcg cac ctg ctc aag cac cat ccc gac ggg			4080
Ala Glu Gln Gly Ser His Ser His Leu Leu Lys His His Pro Asp Gly			
1345	1350	1355	1360

tgc tac gcg cgg atg ctg cag ctg cag cgg ctg acg ggc ggg tgc cgc	4128
Cys Tyr Ala Arg Met Leu Gln Leu Gln Arg Leu Thr Gly Gly Cys Arg	
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gcc cgg gcc gcc gcc gtc gtc gtc caa cgg ggc cgc cgc gta gga tgg	4176
Ala Arg Ala Ala Ala Val Val Val Gln Arg Gly Arg Arg Val Gly Trp	
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Met Asp Gly Ser Trp Met Ser Leu Val Pro	
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35 40 45	
Phe His Leu Ala Ala Gly His Gln Pro Glu Ala Ala Ala Glu Gln Pro	
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Thr Thr Leu Pro Ala Ala Arg Arg Thr Ser Asp Thr Ser Thr Ala Ala	
65 70 75 80	
Gly Ala Ala Pro Pro Ser Pro Ser Pro Pro Pro Pro Ala Pro Leu	
85 90 95	
Glu Met Asp Gln Pro Pro Asn Ala Lys Pro Ala Ser Ser Ser Ala Ala	
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Ala Ala Gly Ala Asn Asp Asn Lys Lys Pro Thr Pro Pro Ala Ala Leu	
115 120 125	
Arg Asp Leu Phe Arg Phe Ala Asp Gly Leu Asp Cys Ala Leu Met Leu	
130 135 140	
Val Gly Thr Leu Gly Ala Leu Val His Gly Cys Ser Leu Pro Val Phe	
145 150 155 160	
Leu Arg Phe Phe Ala Asp Leu Val Asp Ser Phe Gly Ser His Ala Asn	
165 170 175	
Asp Pro Asp Thr Met Val Arg Leu Val Val Lys Tyr Ala Phe Tyr Phe	
180 185 190	
Leu Val Val Gly Ala Ala Ile Trp Ala Ser Ser Trp Ala Glu Ile Ser	

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Cys	Trp	Met	Trp	Thr	Gly	Glu	Arg	Gln	Ser	Thr	Arg	Met	Arg	Ile	Arg
210						215					220				
Tyr	Leu	Asp	Ala	Ala	Leu	Arg	Gln	Asp	Val	Ser	Phe	Phe	Asp	Thr	Asp
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Val	Arg	Thr	Ser	Asp	Val	Ile	Tyr	Ala	Ile	Asn	Ala	Asp	Ala	Val	Val
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Gly	Ala	Gly	Arg	His	Gln	Arg	Glu	Ala	Gly	Gln	Pro	His	Pro	Leu	His
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Gly	His	Leu	Arg	Gly	Gly	Leu	Arg	Arg	Gly	Leu	His	Arg	Arg	Leu	Ala
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Arg	Ala	Val	Gly	Arg	Gln	Arg	His	Arg	Gly	Ala	Gly	Ala	Arg	Ala	Asp
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Thr	Asp	Arg	Ala	Gly	Leu	Arg	Arg	Arg	Gly	Ala	Arg	Asn	Ala	Gly	Val
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Leu	Gly	Gly	Val	Gly	Arg	Arg	Ala	Glu	Asp	Arg	Leu	Pro	Gln	Arg	Leu
	355						360					365			
Arg	Gln	Gly	Ala	Arg	Pro	Arg	Arg	His	Leu	Leu	His	Arg	Leu	Leu	Leu
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Leu	Arg	Pro	Pro	Ala	Leu	Val	Arg	Arg	Thr	Pro	Arg	Pro	Arg	Asn	His
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Thr	Asn	Gly	Gly	Leu	Ala	Ile	Ala	Thr	Met	Phe	Ser	Val	Met	Ile	Gly
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Ser	Arg	Pro	Asp	Val	Pro	Ile	Leu	Arg	Gly	Phe	Ser	Leu	Ser	Val	Pro
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Ala	Gly	Lys	Thr	Ile	Ala	Leu	Val	Gly	Ser	Ser	Gly	Ser	Gly	Lys	Ser



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Thr	Val	Val	Ser	Leu	Leu	Glu	Arg	Phe	Tyr	Asp	Pro	Ser	Ala	Gly	Gln	
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Ile	Leu	Leu	Asp	Gly	His	Asp	Leu	Lys	Ser	Leu	Lys	Leu	Arg	Trp	Leu	
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Gln	Ala	Glu	Met	Glu	Glu	Ala	Ala	Arg	Val	Ala	Asn	Ala	His	Ser	Phe	
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Ile	Val	Lys	Leu	Pro	Asp	Gly	Tyr	Asp	Thr	Gln	Val	Gly	Glu	Arg	Gly	
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Leu	Gln	Leu	Ser	Gly	Gly	Gln	Lys	Gln	Arg	Ile	Ala	Ile	Ala	Arg	Ala	
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Met	Leu	Lys	Asn	Pro	Ala	Ile	Leu	Leu	Leu	Asp	Glu	Ala	Thr	Ser	Ala	
625					630					635					640	
Leu	Asp	Ser	Glu	Ser	Glu	Lys	Leu	Val	Gln	Glu	Ala	Leu	Asp	Arg	Phe	
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Met	Ile	Gly	Arg	Thr	Thr	Leu	Val	Ile	Ala	His	Arg	Met	Ser	Thr	Ile	
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Arg	Lys	Ala	Asp	Val	Val	Ala	Val	Leu	Gln	Gly	Gly	Pro	Val	Ser	Glu	
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Met	Gly	Ala	His	Asp	Glu	Leu	Met	Ala	Lys	Gly	Glu	Asn	Gly	Thr	Tyr	
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Ala	Lys	Phe	Ile	Arg	Met	Gln	Glu	Gln	Ala	His	Glu	Ala	Ala	Phe	Val	
705					710					715					720	
Asn	Ala	Arg	Arg	Ser	Ser	Ala	Arg	Pro	Ser	Ser	Ala	Arg	Asn	Ser	Val	
725					730					735						
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His	Asp	Pro	His	His	His	His	Arg	Thr	Met	Ala	Asp	Lys	Gln	Leu	Ala	
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Phe	Arg	Ala	Gly	Ala	Ser	Ser	Phe	Leu	Arg	Leu	Ala	Arg	Met	Asn	Ser	
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Pro	Glu	Trp	Ala	Tyr	Ala	Leu	Val	Gly	Ser	Leu	Gly	Ser	Met	Val	Cys	

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Val	Tyr	Tyr	Ala	Pro	Asp	Pro	Arg	Tyr	Met	Lys	Arg	Glu	Ile	Ala	Lys						
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	850					855					860										
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Arg	Val	Arg	Glu	Lys	Met	Phe	Ala	Ala	Val	Leu	Arg	Asn	Glu	Ile	Ala						
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Trp	Phe	Asp	Ala	Asp	Glu	Asn	Ala	Ser	Ala	Arg	Val	Ala	Ala	Arg	Leu						
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		995					1000					1005									
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Pro Val Pro Glu Arg	Pro Lys Gly Glu Val	Glu Leu Lys His Val Asp	
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Phe Ser Tyr Pro Ser Arg	Pro Asp Ile Gln Val	Phe Arg Asp Leu Ser	
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Leu Arg Ala Arg Ala Gly	Lys Thr Leu Ala Leu Val	Gly Pro Ser Gly	
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Cys Gly Lys Ser Ser Val	Leu Ala Leu Val Gln Arg	Phe Tyr Glu Pro	
1170	1175	1180	
Thr Ser Gly Arg Val Leu	Leu Asp Gly Lys Asp Val	Arg Lys Tyr Asn	
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Leu Arg Ala Leu Arg Arg	Val Val Ala Val Ala Pro	Gln Glu Pro Phe	
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Leu Phe Ala Ala Ser Ile	His Asp Asn Ile Ala Tyr	Gly Arg Glu Gly	
1220	1225	1230	
Ala Thr Glu Ala Glu Val	Val Glu Ala Ala Thr Gln	Ala Asn Ala His	
1235	1240	1245	
Arg Phe Ile Ala Ala Leu	Pro Glu Gly Tyr Gly Thr	Gln Val Gly Glu	
1250	1255	1260	
Arg Gly Val Gln Leu Ser	Gly Gly Gln Arg Gln Arg	Ile Ala Ile Ala	
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Ala Glu Gln Gly Ser His	Ser His Leu Leu Lys His	His Pro Asp Gly	
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Cys Tyr Ala Arg Met Leu	Gln Leu Gln Arg Leu Thr	Gly Gly Cys Arg	
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Ala Arg Ala Ala Ala Val	Val Val Gln Arg Gly Arg	Arg Val Gly Trp	
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